

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/733,507
DATE: 03/15/2001
TIME: 17:22:12

Input Set : A:\81601-3.pct.seqlist.txt
Output Set: N:\CRF3\03152001\I733507.raw

3 <110> APPLICANT: Agriculture and Agri-Food Canada; The University of Saskatchewan
5 <120> TITLE OF INVENTION: Cyclin Dependant Kinase Inhibitors as Plant Growth
6 Regulators
8 <130> FILE REFERENCE: 81601-3
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/733,507
C--> 11 <141> CURRENT FILING DATE: 2001-03-15
13 <150> PRIOR APPLICATION NUMBER: CA 2,256,121
14 <151> PRIOR FILING DATE: 1998-12-31
16 <160> NUMBER OF SEQ ID NOS: 16
18 <170> SOFTWARE: PatentIn Ver. 2.0
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22 <212> TYPE: DNA
23 <213> ORGANISM: Arabidopsis thaliana
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47 Val Arg Lys Tyr Arg Lys Ala Lys Gly Ile Val Glu Ala Gly Val Ser
48 5 10 15
50 tca acg tat atg cag cta cgg agc cgg aga att gtt tat gtt aga tcg 153
51 Ser Thr Tyr Met Gln Leu Arg Ser Arg Arg Ile Val Tyr Val Arg Ser
52 20 25 30
54 gaa aaa tca agc tct gtc tcc gtc gtc ggt gat aat gga gtt tcg tcg 201
55 Glu Lys Ser Ser Ser Val Ser Val Val Gly Asp Asn Gly Val Ser Ser
56 35 40 45
58 tct tgt agt gga agc aat gaa tat aag aag aaa gaa tta ata cat ctg 249
59 Ser Cys Ser Gly Ser Asn Glu Tyr Lys Lys Lys Glu Leu Ile His Leu
60 50 55 60 65
62 gag gag gaa gat aaa gat ggt gac act gaa acg tcg acg tat cga cgg 297
63 Glu Glu Glu Asp Lys Asp Gly Asp Thr Glu Thr Ser Thr Tyr Arg Arg
64 70 75 80
66 gtg acg aag agg aag ctt ttt gaa aat ctg aga gag gag gag aaa gaa 345
67 Val Thr Lys Arg Lys Leu Phe Glu Asn Leu Arg Glu Glu Glu Lys Glu

ENTERED

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71 Glu Leu Ser Lys Ser Met Glu Asn Tyr Ser Ser Glu Phe Glu Ser Ala
72          100          105          110
74 gtt aaa gaa tcg tta gat tgt tgt tgt agc ggg agg aaa acg atg gag 441
75 Val Lys Glu Ser Leu Asp Cys Cys Cys Ser Gly Arg Lys Thr Met Glu
76          115          120          125
78 gag acg gtg acg gcg gag gag gag gag aag gcg aaa ttg atg acg gag 489
79 Glu Thr Val Thr Ala Glu Glu Glu Glu Lys Ala Lys Leu Met Thr Glu
80 130          135          140          145
82 atg cca acg gaa tcg gaa att gaa gat ttt ttt gtg gaa gct gag aaa 537
83 Met Pro Thr Glu Ser Glu Ile Glu Asp Phe Phe Val Glu Ala Glu Lys
84          150          155          160
86 caa ctc aaa gaa aaa ttc aag aag aag tac aat ttc gat ttc gag aag 585
87 Gln Leu Lys Glu Lys Phe Lys Lys Lys Tyr Asn Phe Asp Phe Glu Lys
88          165          170          175
90 gag aag cca tta gaa gga cgt tac gaa tgg gta aag tta gag 627
91 Glu Lys Pro Leu Glu Gly Arg Tyr Glu Trp Val Lys Leu Glu
92          180          185          190
94 tgaagaagaa gaagaagttt atggtttttt ttttaacttt ttagatttta atatttcagg 687
96 gaataagtta attttatttt gttgatttgg aaatataaga tttgtaggag gaatgttttt 747
98 agaagtacga aattgcacag aaaaagaaga aagcttttta acagatttta gagccagaa 807
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108 <213> ORGANISM: Arabidopsis thaliana
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118 35 40 45
120 Ser Ser Cys Ser Gly Ser Asn Glu Tyr Lys Lys Lys Glu Leu Ile His
121 50 55 60
123 Leu Glu Glu Glu Asp Lys Asp Gly Asp Thr Glu Thr Ser Thr Tyr Arg
124 65 70 75 80
126 Arg Val Thr Lys Arg Lys Leu Phe Glu Asn Leu Arg Glu Glu Glu Lys
127 85 90 95
129 Glu Glu Leu Ser Lys Ser Met Glu Asn Tyr Ser Ser Glu Phe Glu Ser
130 100 105 110
132 Ala Val Lys Glu Ser Leu Asp Cys Cys Cys Ser Gly Arg Lys Thr Met
133 115 120 125
135 Glu Glu Thr Val Thr Ala Glu Glu Glu Glu Lys Ala Lys Leu Met Thr
136 130 135 140
138 Glu Met Pro Thr Glu Ser Glu Ile Glu Asp Phe Phe Val Glu Ala Glu
139 145 150 155 160

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141 Lys Gln Leu Lys Glu Lys Phe Lys Lys Lys Tyr Asn Phe Asp Phe Glu
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158 aagaaagaat taatacatct ggaggaggaa gataaagatg gtgacactga aacgtcgacg 180
160 tatcgacggg gtacgaagag gaagcttttt gaaaatctga gagaggagga gaaagaagaa 240
162 ttaagtaaat ccatggagaa ttattcatcg gaatttgaat cggcggttaa agaatcgtta 300
164 gattgttggt gtagcgggag gaaaacgatg gaggagacgg tgacggcgga ggaggaggag 360
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168 gctgagaaac aactcaaaga aaaattcaag aagaagtaca atttcgattt cgagaaggag 480
170 aagccattag aaggacgtta cgaatgggta aagtttagag gaagaagaag aagaagttta 540
172 tggttttttt ttttaacttt tagattttta tatttcaggg aataagttta ttttattttg 600
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180 <213> ORGANISM: Arabidopsis thaliana
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187 atcgttagat tgttggttga gcgggaggaa aacgatggag gagacggtga cggcggagga 180
189 ggaggagaag gcgaaattga tgacggagat gccaacgga tcggaaattg aagatttttt 240
191 tgtggaagct gagaaacaac tcaaagaaaa attcaagaag aagtacaatt tcgatttcga 300
193 gaaggagaag ccattagaag gacgttacga atgggtaaag ttagagtga gaagaagaag 360
195 aagtttatgg tttttttttt aacttttttag attttaatat ttcagggaat aagttaattt 420
197 tattttggtg atttggaat ata                                     443
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203 <213> ORGANISM: Arabidopsis thaliana
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210 gtttagattgt tgttgtagcg ggaggaaaac gatggaggag gaggaggaga aggcgaaatt 180
212 gatgacggag atgccaacgg aatcggaat tgaagatttt tttgtggaag ctgagaaaca 240
214 actcaaagaa aaattcaaga agaagtacaa ttctgatttc gagaaggaga agccattaga 300
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222 <211> LENGTH: 755
223 <212> TYPE: DNA
224 <213> ORGANISM: Arabidopsis thaliana

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231 cctccggttg aagaacagtg tcaaatcgaa gaagaagatt cgtcggttct gtgttgttct 180
233 acatcggaag agaaatcgaa acggagaatc gaattttagt atcttgagga aaataacggt 240
235 gacgatcggt aaacagaaac gtcgtggatt tacgatgatt tgaataagag tgaggaaatcg 300
237 atgaacatgg attcttcttc ggtggctgtt gaagatgtag agtctcgccg cagggttaagg 360
239 aagagtctcc atgagacggt gaaggaagct gagttagaag acttttttca ggtggcgagg 420
241 aaagatcttc ggaataagtt gttggaatgt tctatgaagt ataacttoga ttctgagaaa 480
243 gatgagccac ttggtggagg aagatacgag tgggttaaat tgaatccatg aagaagacga 540
245 tgatgataat gatgatcatt gttttcacca aagtacttat tatttctctt ctgtaataat 600
247 ctttgccttg atttttcttt taacaaaatc caaatgtaga tatctttctc tcgaataatc 660
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264 cttaatgatt cttctgattc atcaccggac tctcatgacg tcatcgtctt cgcggtttca 180
266 tcttcttccg ttgcttcgct ggcggcttta gcgtctgatg aatgttccgt taccatcggt 240
268 ggagaagaaa gtgatcagtc ctcgagtatc agctccggtt gtttcaccag tgaatcgaaa 300
270 gaaatcgcga agaacagttc gtcgttttgt gtagatctgg aggatcatca aatcgaaaac 360
272 gaaaccgaaa cctcaacatt catcaccagc aatttcagaa aagagacgag tccagtgagt 420
274 gagggttttg gagaaacgac aacagaaatg gaatcatcat cggcaacgaa gagaaaacaa 480
276 cggggggtga ggaagactcc aacggcgggc gagattgagg atttgttctc ggagctagag 540
278 agtcaagacg ataagaagaa gcaattcata gaaaagtaca acttcgatat tgtcaatgac 600
280 gaaccgcttg aaggctcgct caagtgggat cgactttaag ccatcaaaaa gcaaatacca 660
282 tccatgaaga agacaaaaga aaaatagggt ttgtttttcg tggttaacat ttccacttgt 720
284 acagctctag tctatttctc tttaaaaacc tatgttacta gttcgtacaa aacaaaacaa 780
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299 ttacgctcac cgacgagaac tctttcttct tcttctctt cttctctggc ttactcggtt 180
301 tcagattccg gaggtttctg ctccgtcgcg ttatctgaag aagaagacga tcatctaagc 240
303 tcaagcatca gctctggttg ttccagcagc gaaactaacg aaatcgctac tcgtcttcca 300
305 ttttcagatc tggaggctca tgaaatctcc gaaaccgaaa tctcaacggt actcaccaac 360
307 aatttcagga aacagggaat ttcatcaagc gagaatctgg gagaacagc agaaatggac 420
309 tcggcgacga cggagatgag agatcagaga aagacggaga agaagaagaa gatggaaaaa 480
311 tcaccgacgc aggcagagct tgatgacttt ttctcggcgg cggagagata cgaacagaaa 540
313 cgattcacag aaaagtacaa ctacgacatc gtcaatgata cgccgcttga aggtcggtac 600
315 cagtgggtta gtctgaaacc ttagaagcca tggaagaaca aa 642

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328 aacagtctcg gttttgaatc aagacacagc acaagggaga gcacgccttg taactttgtt 180
330 gaggatatgg agatcatggt tacaccaggg tctagcacga ggtcgatgtg cagagcaacc 240
332 aaagagtaca caagggaaca agataacgtg atccccacca ctagtgaat ggaggagttc 300
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336 gtgaatgata tccccctcag cggacgttac gaatgggtgc aagtcaaacc atgaagttca 420
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356 35 40 45
358 Ser Ser Cys Ser Gly Ser Asn Glu Tyr Lys Lys Lys Glu Leu Ile His
359 50 55 60
361 Leu Glu Glu Glu Asp Lys Asp Gly Asp Thr Glu Thr Ser Thr Tyr Arg
362 65 70 75 80
364 Arg Gly Thr Lys Arg Lys Leu Cys Glu Asn Leu Arg Glu Glu Glu Lys
365 85 90 95
367 Glu Glu Leu Ser Lys Ser Met Glu Asn Tyr Ser Ser Glu Phe Glu Ser
368 100 105 110
370 Ala Val Lys Glu Ser Leu Asp Cys Cys Cys Ser Gly Arg Lys Thr Met
371 115 120 125
373 Glu Glu Thr Val Thr Ala Glu Glu Glu Glu Lys Ala Lys Leu Met Thr
374 130 135 140
376 Glu Met Pro Thr Glu Ser Glu Ile Glu Asp Phe Phe Val Glu Ala Glu
377 145 150 155 160
379 Lys Gln Leu Lys Glu Lys Phe Lys Lys Lys Tyr Asn Phe Asp Phe Glu
380 165 170 175
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VERIFICATION SUMMARY

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L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

STATISTICS SUMMARY

PATENT APPLICATION: US/09/733,507

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Input Set : A:\81601-3.pct.seqlist.txt

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Application Serial Number: US/09/733,507

Alpha or Numeric: Numeric

Application Class:

Application File Date: 03-15-2001

Art Unit: OIPE

Software Application: PatentIn

Total Number of Sequences: 16

Total Nucleotides: 5942

Total Amino Acids: 1311

Number of Errors: 0

Number of Warnings: 0

Number of Corrections: 2

MESSAGE SUMMARY

270 C: 1 (Current Application Number differs)

271 C: 1 (Current Filing Date differs)